Pham 107048

1: Nairb_35 + 3

2: RonRayGun_36 + 1

3: Mendokysei_34

Note: Tracks are now grouped by subcluster and scaled. Switching in subcluster is indicated by changes in track color. Track scale is now set by default to display the region 30 bp upstream of start 1 to 30 bp downstream of the last possible start. If this default region is judged to be packed too tightly with annotated starts, the track will be further scaled to only show that region of the ORF with annotated starts. This action will be indicated by adding "Zoomed" to the title. For starts, yellow indicates the location of called starts comprised solely of Glimmer/GeneMark auto-annotations, green indicates the location of called starts with at least 1 manual gene annotation.

Pham 107048 Report

This analysis was run 04/05/24 on database version 557.

Pham number 107048 has 7 members, 0 are drafts.

Phages represented in each track:
• Track 1 : Nairb_35, Ibrahim_36, Whitty_36, Bernal13_36
• Track 2 : RonRayGun_36, ZenTime222_35
• Track 3 : Mendokysei_34

Summary of Final Annotations (See graph section above for start numbers):

The start number called the most often in the published annotations is 1, it was called in 5 of the 7 non-draft genes in the pham.

Genes that call this "Most Annotated" start:
• Bernal13_36, Ibrahim_36, Mendokysei_34, Nairb_35, Whitty_36,

Genes that have the "Most Annotated" start but do not call it:
• RonRayGun_36, ZenTime222_35,

Genes that do not have the "Most Annotated" start:
•

Summary by start number:

Start 1:
• Found in 7 of 7 (100.0%) of genes in pham
• Manual Annotations of this start: 5 of 7
• Called 71.4% of time when present
• Phage (with cluster) where this start called: Bernal13_36 (T), Ibrahim_36 (T), Mendokysei_34 (T), Nairb_35 (T), Whitty_36 (T),

Start 3:
• Found in 7 of 7 (100.0%) of genes in pham
• Manual Annotations of this start: 2 of 7
• Called 28.6% of time when present
• Phage (with cluster) where this start called: RonRayGun_36 (T), ZenTime222_35 (T),

Summary by clusters:

There is one cluster represented in this pham: T

Info for manual annotations of cluster T:
•Start number 1 was manually annotated 5 times for cluster T.
•Start number 3 was manually annotated 2 times for cluster T.

Gene Information:
Gene: Bernal13_36 Start: 30000, Stop: 30314, Start Num: 1
Candidate Starts for Bernal13_36:
(Start: 1 @30000 has 5 MA's), (Start: 3 @30036 has 2 MA's), (4, 30054), (5, 30162), (6, 30186), (7, 30249),

Gene: Ibrahim_36 Start: 30446, Stop: 30760, Start Num: 1
Candidate Starts for Ibrahim_36:
(Start: 1 @30446 has 5 MA's), (Start: 3 @30482 has 2 MA's), (4, 30500), (5, 30608), (6, 30632), (7, 30695),

Gene: Mendokysei_34 Start: 29535, Stop: 29852, Start Num: 1
Candidate Starts for Mendokysei_34:
(Start: 1 @29535 has 5 MA's), (2, 29562), (Start: 3 @29571 has 2 MA's), (4, 29589), (5, 29700), (7, 29787),

Gene: Nairb_35 Start: 30000, Stop: 30314, Start Num: 1
Candidate Starts for Nairb_35:
(Start: 1 @30000 has 5 MA's), (Start: 3 @30036 has 2 MA's), (4, 30054), (5, 30162), (6, 30186), (7, 30249),

Gene: RonRayGun_36 Start: 30482, Stop: 30760, Start Num: 3
Candidate Starts for RonRayGun_36:
(Start: 1 @30446 has 5 MA's), (Start: 3 @30482 has 2 MA's), (4, 30500), (5, 30608), (6, 30632), (7, 30695),

Gene: Whitty_36 Start: 30000, Stop: 30314, Start Num: 1
Candidate Starts for Whitty_36:
(Start: 1 @30000 has 5 MA's), (Start: 3 @30036 has 2 MA's), (4, 30054), (5, 30162), (6, 30186), (7, 30249),

Gene: ZenTime222_35 Start: 30036, Stop: 30314, Start Num: 3
Candidate Starts for ZenTime222_35:
(Start: 1 @30000 has 5 MA's), (Start: 3 @30036 has 2 MA's), (4, 30054), (5, 30162), (6, 30186), (7, 30249),